

logo

NTDB id 1149749 DXE47 RS12705 WP 000287260.1
NTDB id 39 MW RS04875 WP 000287265.1
NTDB id 23 SA RS04995 WP 000287265.1
consensus

MYSQNIYVIRKGMVIRPAFDDDDQRRNGSEIRFDKTRIQNPFKVKIERSCKFYGNTYLGKKAETNRTIGISSKPPIL
 MYSQNIYVIRKGMVIRPAFDDDDQRRNGSEIRFDKTRIQNPFKVKIERSCKFYGNTYLGKKAETNRTIGISSKPPIL
 MYSQNIYVIRKGMVIRPAFDDDDQRRNGSEIRFDKTRIQNPFKVKIERSCKFYGNTYLGKKAETNRTIGISSKPPIL
 MYSQNIYVIRKGMVIRPAFDDDDQRRNGSEIRFDKTRIQNPFKVKIERSCKFYGNTYLGKKAETNRTIGISSKPPIL
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80
80
80

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NTDB id 23 SA RS04995 WP 000287265.1
consensus

L T P L F P T Y F F P T H S D R Q N E N I W L N M H Y I E S I K E L K N R K C K V T F I N N E S I I L H V S Y H S L W H Q Y N N S I F Y Y Y M V D K Q S R M I S
 L T P L F P T Y F F P T H S D R Q N E N I W L N M H Y I E S I K E L K N R K C K V T F I N N E S I I L H V S Y H S L W H Q Y N N S I F Y Y Y M V D K Q S R M I S
 L T P L F P T Y F F P T H S D R Q N E N I W L N M H Y I E S I K E L K N R K C K V T F I N N E S I I L H V S Y H S L W H Q Y N N S I F Y Y Y M V D K Q S R M I S
 L T P L F P T Y F F P T H S D R Q N E N I W L N M H Y I E S I K E L K N R K C K V T F I N N E S I I L H V S Y H S L W H Q Y N N S I F Y Y Y M V D K Q S R M I S

160
160
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Sequence logo for the 16-mer motif KNPDQPIDYNKATLNVFEALTRYSLFEDK. The logo shows the conservation of each amino acid at each position. The sequence is repeated four times below the logo.

189
189
189

-  non conserved
-  similar
-  > 50% conserved